

55084US-sequencelisting.ST25
SEQUENCE LISTING

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ARTYMIUK, Peter
SAYERS, Jon

<120> Polypeptide Variants

<130> 100042.55084US

<140> TBA
<141> 2004-07-23

<150> PCT/GB03/00253
<151> 2003-01-24

<150> GB 0201679.8
<151> 2003-01-24

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<170> PatentIn version 3.2

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Gly Gly Gly Gly Ser
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aagtattcat tcctgcagaa cccccagacc tccctctgtt tctcagagtc tattccgaca 180
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Phe Pro Thr Ile Pro Leu Ser Arg Leu Phe Asp Asn Ala Ser Leu Arg
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Ala His Arg Leu His Gln Leu Ala Phe Asp Thr Tyr Gln Glu Phe Glu
20 25 30

Glu Ala Tyr Ile Pro Lys Glu Gln Lys Tyr Ser Phe Leu Gln Asn Pro
35 40 45

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Gln Thr Ser Leu Cys Phe Ser Glu Ser Ile Pro Thr Pro Ser Asn Arg
50 55 60

Glu Glu Thr Gln Gln Lys Ser Asn Leu Glu Leu Leu Arg Ile Ser Leu
65 70 75 80

Leu Leu Ile Gln Ser Trp Leu Glu Pro Val Gln Phe Leu Arg Ser Val
85 90 95

Phe Ala Asn Ser Leu Val Tyr Gly Ala Ser Asp Ser Asn Val Tyr Asp
100 105 110

Leu Leu Lys Asp Leu Glu Glu Gly Ile Gln Thr Leu Met Gly Arg Leu
115 120 125

Glu Asp Gly Ser Pro Arg Thr Gly Gln Ile Phe Lys Gln Thr Tyr Ser
130 135 140

Lys Phe Asp Thr Asn Ser His Asn Asp Asp Ala Leu Leu Lys Asn Tyr
145 150 155 160

Gly Leu Leu Tyr Cys Phe Arg Lys Asp Met Asp Lys Val Glu Thr Phe
165 170 175

Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
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taccaacaga tcctcaccag tatgccttcc agaaacgtga tccaaatatc caacgacctg 240
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acagaggtgg tggccctgag caggctgcag gggctctctgc aggacatgct gtggcagctg 420
 gacctcagcc ctgggtgc 438

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 ttatttcctg ggctccaccc catcctgacc ttatccaaga tggaccagac actggcagtc 180
 taccaacaga tcctcaccag tatgccttcc agaaacgtga tccaaatata caacgacctg 240
 gagaacctcc gggatcttct tcacgtgctg gccttctcta agagctgcca cttgccctgg 300
 gccagtggcc tggagacctt ggacagcctg ggggggtgtcc tggaagcttc aggctactcc 360
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Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr Gln Ser Val Ser Ser
 20 25 30

Lys Gln Lys Val Thr Gly Leu Asp Phe Ile Pro Gly Leu His Pro Ile
 35 40 45

Leu Thr Leu Ser Lys Met Asp Gln Thr Leu Ala Val Tyr Gln Gln Ile
 50 55 60

Leu Thr Ser Met Pro Ser Arg Asn Val Ile Gln Ile Ser Asn Asp Leu
 65 70 75 80

Glu Asn Leu Arg Asp Leu Leu His Val Leu Ala Phe Ser Lys Ser Cys
 85 90 95

His Leu Pro Trp Ala Ser Gly Leu Glu Thr Leu Asp Ser Leu Gly Gly
 100 105 110

Val Leu Glu Ala Ser Gly Tyr Ser Thr Glu Val Val Ala Leu Ser Arg
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Leu Gln Gly Ser Leu Gln Asp Met Leu Trp Gln Leu Asp Leu Ser Pro
130 135 140

Gly Cys
145

